



Fig. 1

HRGP	173	ASFRVDR	IER-V-ARVRGG--EGT-YFVDFSVR-----NCPRHHFPRHPNVFGFCRAD--LEFYDVEALDL
CD36	89	GPYTYR	VR--FLAKENVTDDEA-DNTVSE-LQP-----NGA--IFEPSLSV-GT-EADN-ETVLNLAVA
LIMP-II	90	GPYTYR	E--LRNKANIQFGDNG-TTIS-AVT-----NKA-YIFERNQSV-GDPKID-LIRTLNIPVL
HRGP	443	GPREFCRQIGSVVRLPPLRKGEVLPLPEA-NFPSEPLRHHKHLKPDNQP----FPQSVSESCPGKFKSGEPQVSMFT	
CD36	89	GPYTYR	FLAKENVTDDEA-DNTVSE-LQP-----NGA--IFEPSLSVGT--EADN-ETVLNLAVA
LIMP-II	90	GPYTYR	E-LRNKANIQFG-DNGTTIS-AVT-----NKAY-IFERNQSVGDP-KID-LIRTLNIPVL



Fig. 2

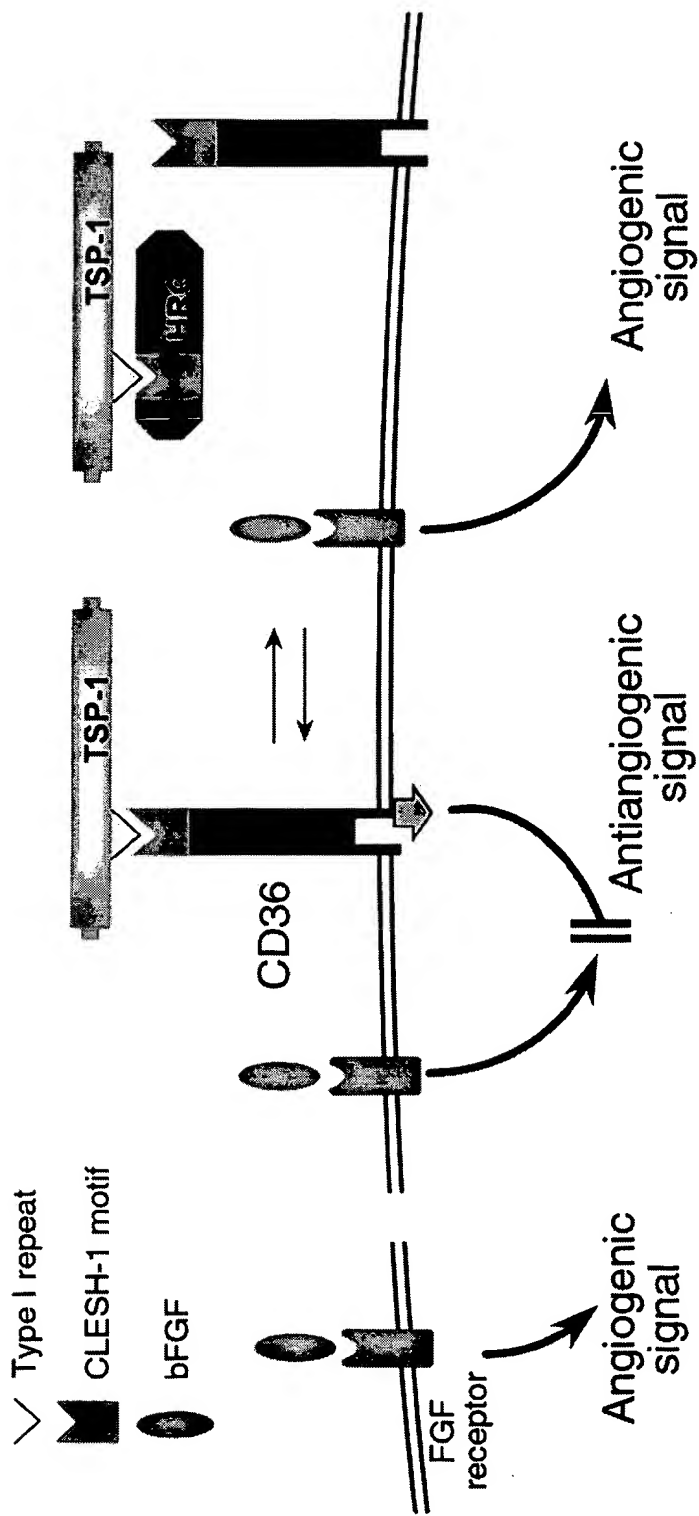




Fig. 3

